

SEQUENCE LISTING

<110> Biomolecular Engineering Research Institute

<120> Crystal of Protein Complex, Structural Coordinate Thereof and Use Thereof

<130> 661699

<150> JP 56905/1999 and 218691/1999

<151> 1999-03-04 and 1999-08-02

<160> 3

<210> 1

<211> 175

<212> PRT

<213> homo sapience

<400> 1

Met Thr Pro Leu Gly Pro Ala Ser Ser Leu Pro Gln Ser Phe Leu Leu

1 5 10 15

Lys Cys Leu Glu Gln Val Arg Lys Ile Gln Gly Asp Gly Ala Ala Leu

20 25 30

Gln Glu Lys Leu Cys Ala Thr Tyr Lys Leu Cys His Pro Glu Glu Leu

35 40 45

Val Leu Leu Gly His Ser Leu Gly Ile Pro Trp Ala Pro Leu Ser Ser

50 55 60

Cys Pro Ser Gln Ala Leu Gln Leu Ala Gly Cys Leu Ser Gln Leu His

65 70 75 80

00449 09130

Ser Gly Leu Phe Leu Tyr Gln Gly Leu Leu Gln Ala Leu Glu Gly Ile

85

90

95

Ser Pro Glu Leu Gly Pro Thr Leu Asp Thr Leu Gln Leu Asp Val Ala

100

105

110

Asp Phe Ala Thr Thr Ile Trp Gln Gln Met Glu Glu Leu Gly Met Ala

115

120

125

Pro Ala Leu Gln Pro Thr Gln Gly Ala Met Pro Ala Phe Ala Ser Ala

130

135

140

Phe Gln Arg Arg Ala Gly Gly Val Leu Val Ala Ser His Leu Gln Ser

145

150

155

160

Phe Leu Glu Val Ser Tyr Arg Val Leu Arg His Leu Ala Gln Pro

165

170

175

<210> 2

<211> 215

<212> PRT

<213> mouse

<400> 2

Ala Gly Tyr Pro Pro Ala Ser Pro Ser Asn Leu Ser Cys Leu Met His

1

5

10

15

Leu Thr Thr Asn Ser Leu Val Cys Gln Trp Glu Pro Gly Pro Glu Thr

20

25

30

His Leu Pro Thr Ser Phe Ile Leu Lys Ser Phe Arg Ser Arg Ala Asp

35

40

45

Cys Gln Tyr Gln Gly Asp Thr Ile Pro Asp Cys Val Ala Lys Lys Arg

50

55

60

Gln Asn Asn Cys Ser Ile Pro Arg Lys Asn Leu Leu Leu Tyr Gln Tyr

65

70

75

80

00044619 091601

His Leu Pro Thr Ser Phe Thr Leu Lys Ser Phe Lys Ser Arg Gly Asn

35	40	45
Cys Gln Thr Gln Gly Asp Ser Ile Leu Asp Cys Val Pro Lys Asp Gly		
50	55	60
Gln Ser His Cys Cys Ile Pro Arg Lys His Leu Leu Leu Tyr Gln Asn		
65	70	75
Met Gly Ile Trp Val Gln Ala Glu Asn Ala Leu Gly Thr Ser Met Ser		
85	90	95
Pro Gln Leu Cys Leu Asp Pro Met Asp Val Val Lys Leu Glu Pro Pro		
100	105	110
Met Leu Arg Thr Met Asp Pro Ser Pro Glu Ala Ala Pro Pro Gln Ala		
115	120	125
Gly Cys Leu Gln Leu Cys Trp Glu Pro Trp Gln Pro Gly Leu His Ile		
130	135	140
Asn Gln Lys Cys Glu Leu Arg His Lys Pro Gln Arg Gly Glu Ala Ser		
145	150	155
Trp Ala Leu Val Gly Pro Leu Pro Leu Glu Ala Leu Gln Tyr Glu Leu		
165	170	175
Cys Gly Leu Leu Pro Ala Thr Ala Tyr Thr Leu Gln Ile Arg Cys Ile		
180	185	190
Arg Trp Pro Leu Pro Gly His Trp Ser Asp Trp Ser Pro Ser Leu Glu		
195	200	205
Leu Arg Thr Thr Glu Arg Ala		
210	215	

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